

SEQUENCE LISTING

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360

420

480

505

<120> Adeno-associated virus vectors <130> 875.007US2 <140> US10/054,665 <141> 2002-01-22 <150> US 60/086,166 <151> 1998-05-20 <150> US 09/276,625 <151> 1999-03-25 <160> 14 <170> FastSEQ for Windows Version 4.0 <210> 1 <211> 20 <212> DNA <213> Adeno-associated virus 400> 1 cgggggtcgt tgggcggtca 20 <210> 2 <211> 19 <212> DNA <213> Adeno-associated virus <400> 2 gggcggagcc tatggaaaa 19 <210> 3 <211> 505 <212> DNA <213> Artificial Sequence <223> A synthetic consensus sequence <400> 3 cgggggtcgt tgggcggtca gccaggcggg ccatttaccg taagttatgt aacgactgca 60 ggcatgcaag ctcgaattca tcggtagata agtagcatgg cgggttaatc attaactaca 120 aggaacccct agtgatggag ttggccactc cctctctgcg cgctcgctcg ctcgctgagg 180 ccgggcgacc aaaggtcgcc cgacgcccgg gctttgcccg ggcggcctca gtgagcgagc 240 gagegegeag etgegegete getegeteae tgaggeegee egggeaaage eegggegteg 300 ggcgaccttt ggtcgcccgg cctcagcgag cgagcgagcg cgcagagagg gagtggccaa

ctccatcact aggggttcct tgtagttaat gattaacccg ccatgctact tatctacagc

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tggcgttttt ccataggctc cgccc

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<211> 272
<212> DNA
<213> AAV circular intermediate, clone p81
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                                                                         60
tgatggagtt ggccactccc tctctgcgcg ctcgctcgct cactgaggcc gggcggccaa
                                                                        120
aggtcgcccg acgcccgggc tttgcccggg cggcctcagt gagcgagcga gcgcgcagag
                                                                        180
agggagtggc caactccatc actaggggtt ccttgtagtt aatgattaac ccgccatgct
                                                                        240
acttatctac cgatgaattc gagcttgcat gc
                                                                        272
<210> 5
<211> 300
<212> DNA
<213> AAV circular intermediate, clone p79
<400> 5
gcatgcaagc tgtagataag tagcatggcg ggttaatcat taactacaag gaacccctag
                                                                         60
tgatggagtt ggccactccc tetetgegeg etegeteget caetgaggee gggegegege
                                                                        120
tegetegete actgaggeeg ggegaceaaa ggtegeeega geeegggett tgeeegggeg
                                                                        180
gcctcagtga gcgagcgcgc gcgcagagag ggagtggcca actccatcac taggggttcc
                                                                        240
ttgtagttaa tgattaaccc gccatgctac ttatctaccg atgaattcga gcttgcatgc
                                                                        300
<210> 6
<211> 272
<212> DNA
<213> AAV circular intermediate, clone p1202
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                                                                         60
tgatggagtt ggccactccc tctctgcgcg ctcgctcgct cactgaggcc gggcgaccaa
                                                                        120
aggtcgcccg acgcccgggc tttggtcgcc cggcctcagt gagcgagcga gcgcgcagag
                                                                        180
agggagtggc caactccatc actaggggtt ccttgtagtt aatgattaac ccgccatgct
                                                                       240
acttatctac cgatgaattc gagcttgcat gc
                                                                       272
<210> 7
<211> 165
<212> DNA
<213> Unknown
<220>
<223> SEQ ID NO:1 of U.S. Patent No. 5,478,745
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ccgggcgacc aaaggtcgcc cgacgcccgg gctttgcccg ggcggcctca gtgagcgagc
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gagcgcgcag agagggagtg gccaactcca tcactagggg ttcct
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<210> 8
<211> 282
<212> DNA
<213> rAAV circular intermediate, clone p79
<400> 8
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tagataagta gcatggcggg ttaatcattg cctacaaaga gcccctagtg atggagtggg
                                                                       120
ccactccctc tcttcgccga gcgcgcagag agggagtggc caactccctc actaggggtt
                                                                       180
cctggcagtt aatgattaac ccgccatgct acttatctac agcttgcatg catgtgagca
                                                                       240
aaaggccagc aaaaggccag gaaccgtaaa aaggccgcgt tg
                                                                       282
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₹211> 345
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<213> rAAV circular intermediate, clone p80
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taagtagcat ggcgggttaa tcattaacta caaggaaccc ctagtgatgg agttggccac
                                                                        120
tecetetetg egegeteget egetegetea ggeegggega ecaaaggteg eeegaegeee
                                                                        180
gcccggcctc agcgagcgag cgagcgcgca gagagggagt ggccaactcc atcactaggg
                                                                        240
gttccttgta gttaatgatt aacccgccat gctacttatc tacagcttgc atgcatgtga
                                                                        300
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                                                                        345
<210> 10
<211> 276
<212> DNA
<213> rAAV circular intermediate, clone p81
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taagtagcat ggcgggttaa tcattgccta caaagagccc ctagtgatgg agcccggcct
                                                                        120
caccgagcga gcgagcgcgc agagagggag tggccaactc catcactagg ggttccttgt
                                                                        180
agttaatgat taacccgcca tgctacttat ctacagcttg catgcatgtg agcaaaaggc
                                                                        240
cagcaaaagg ccaggaaccg taaaaaggcc gcgttg
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<210> 11
<211> 316
<212> DNA
<213> rAAV circular intermediate, clone p86
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taagtagcat ggcgggttaa tcattaacta caaggaaccc ctagtgatgg agttggccac
                                                                        120
tecetetetg egegeteget egetegetga ggeegeeeeg geeteagega gegagegage
                                                                        180
gegeagagag ggaetggeea actecateae taggggttee ttgtagttaa tgattaacce
                                                                        240
gccatgctac ttatctacag cttgcatgca tgtgagcaaa aggccagcaa aaggccagga
                                                                        300
accgtaaaaa ggccgc
                                                                       316
<210> 12
<211> 208
<212> DNA
<213> rAAV circular intermediate, clone p87
<400> 12
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taagtagcat ggcgggttac tcattgccta caaagagccc ctagtgatgg aattggaatg
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attcaccctc catgctactt atctacagct tgcatgcatg tgagcaaaag gccagcaaaa
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ggccaggaac cgtaaaaagg ccgcgttg
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<210> 13
<211> 310
<212> DNA
<213> rAAV circular intermediate, clone p88
<400> 13
gccatttacc gtaagttatg taacgactgc aggcatgcaa gctcgaattc atcggtagat
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aagtagcatg gcgggttaat cattgcctac aaagagcccc tagtgatgga gttggccact
                                                                       120
ccctctctgc gcgctcgctc gctgggcccg gcctcagcga gcgagcgagc gcgcagagag
                                                                       180
ggagtggcca actccatcac taggggttcc ttgtagttaa tgattaaccc gccatgctac
                                                                       240
ttatctacag cttgcatgca tgtgagcaaa aggccagcaa aaggccagga accgtaaaaa
                                                                       300
ggccgcgttg
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4,

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<210> '14
<211> 334
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<213> Artificial Sequence
<220>
<223> A synthetic portion of the consensus sequence
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                                                                       120
cgcccgggct ttgcccgggc ggcctcagtg agcgagcgag cgcgcagctg cgcgctcgct
                                                                       180
cgctcactga ggccgcccgg gcaaagcccg ggcgtcgggc gacctttggt cgcccggcct
                                                                       240
cagcgagcga gcgagcgcgc agagagggag tggccaactc catcactagg ggttccttgt
                                                                       300
agttaatgat taacccgcca tgctacttat ctac
                                                                      334
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